

0260

OIPE

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RAW SEQUENCE LISTING PATENT APPLICATION US/09/459,774

DATE: 12/30/1999
TIME: 14:42:06

Input Set: I459774.RAW

This Raw Listing contains the General Information
Section and up to first 5 pages.

ENTERED

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1  <110> APPLICANT: Michael Robert Barnes
2      Tania Tamson Testa
3  <120> TITLE OF INVENTION: NOVEL COMPOUNDS
4  <130> FILE REFERENCE: GP-30193
5  <140> CURRENT APPLICATION NUMBER: US/09/459,774
6  <141> CURRENT FILING DATE: 1999-12-13
7  <150> EARLIER APPLICATION NUMBER: UK 9828419.3
8  <151> EARLIER FILING DATE: 1998-12-23
9  <160> NUMBER OF SEQ ID NOS: 2
10 <170> SOFTWARE: FastSEQ for Windows Version 3.0
11 <210> SEQ ID NO 1
12 <211> LENGTH: 1050
13 <212> TYPE: DNA
14 <213> ORGANISM: HOMO SAPIENS
15 <400> SEQUENCE: 1
16      atgaaccgga aagcgcggcg ctgcctgggc cacctctttc tcagcctggg catggtctac      60
17      ctccggatcg gtggcttctc ctcagtggta gctctgggcg caagcatcat ctgtaacaag      120
18      atcccaggcc tggctccagc acagcgggcg atctgccaga gccggcccga cgccatcatc      180
19      gtcataaggag aaggctcaca aatgggcctg gacgagtgtc agtttcagtt ccgcaatggc      240
20      cgctggaact gctctgcact gggagagcgc accgtcttcg ggaaggagct caaagtgggg      300
21      agccgggagg ctgcgttcac ctacgccatc attgcccgcg gcgtggccca cgccatcaca      360
22      gctgcctgta cccaggggcaa cctgagcgac tgtggctgcg acaaagagaa gcaaggccag      420
23      taccaccggg acgagggctg gaagtggggt ggctgctctg ccgacatccg ctacggcatc      480
24      ggcttcgcca aggtctttgt ggatgcccgg gagatcaagc agaatggccg gactctcatg      540
25      aacttgcaca acaacgaggc aggccgaaag atcctggagg agaacatgaa gctgggaatgt      600
26      aagtgccacg gcgtgtcagg ctcggtgcacc accaagacgt gctggaccac actgccacag      660
27      ttctgggagc tgggctacgt gctcaaggac aagtacaacg aggccgttca cgtggagcct      720
28      gtgcgtgcca gccgcaacaa gccggccacc ttctgaaga tcaagaagcc actgtcgtac      780
29      cgcaagccca tggacacgga cctgggtgtac atcgagaagt cgcccaacta ctgcgaggag      840
30      gacccggtga ccggcagtgt gggcacccag ggccgcgcct gcaacaagac ggctccccag      900
31      gccagcggct gtgacctcat gtgctgtggg cgtggctaca acacccacca gtacggccgc      960
32      gtgtggcagt gcaactgtaa gttccactgg tgctgctatg tcaagtgcaa cacgtgcagc      1020
33      gagcgcacgg agatgtacac gtgcaagtga      1050
34 <210> SEQ ID NO 2
35 <211> LENGTH: 349
36 <212> TYPE: PRT
37 <213> ORGANISM: HOMO SAPIENS
38 <400> SEQUENCE: 2
39      Met Asn Arg Lys Ala Arg Arg Cys Leu Gly His Leu Phe Leu Ser Leu
40      1          5          10          15
41      Gly Met Val Tyr Leu Arg Ile Gly Gly Phe Ser Ser Val Val Ala Leu
42      20          25          30
43      Gly Ala Ser Ile Ile Cys Asn Lys Ile Pro Gly Leu Ala Pro Arg Gln
44      35          40          45

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45   Arg Ala Ile Cys Gln Ser Arg Pro Asp Ala Ile Ile Val Ile Gly Glu
46       50                               55                               60
47   Gly Ser Gln Met Gly Leu Asp Glu Cys Gln Phe Gln Phe Arg Asn Gly
48   65                               70                               75                               80
49   Arg Trp Asn Cys Ser Ala Leu Gly Glu Arg Thr Val Phe Gly Lys Glu
50                               85                               90                               95
51   Leu Lys Val Gly Ser Arg Glu Ala Ala Phe Thr Tyr Ala Ile Ile Ala
52                               100                              105                              110
53   Ala Gly Val Ala His Ala Ile Thr Ala Ala Cys Thr Gln Gly Asn Leu
54   115                              120                              125
55   Ser Asp Cys Gly Cys Asp Lys Glu Lys Gln Gly Gln Tyr His Arg Asp
56   130                              135                              140
57   Glu Gly Trp Lys Trp Gly Gly Cys Ser Ala Asp Ile Arg Tyr Gly Ile
58   145                              150                              155                              160
59   Gly Phe Ala Lys Val Phe Val Asp Ala Arg Glu Ile Lys Gln Asn Ala
60                               165                               170                               175
61   Arg Thr Leu Met Asn Leu His Asn Asn Glu Ala Gly Arg Lys Ile Leu
62                               180                               185                               190
63   Glu Glu Asn Met Lys Leu Glu Cys Lys Cys His Gly Val Ser Gly Ser
64   195                              200                              205
65   Cys Thr Thr Lys Thr Cys Trp Thr Thr Leu Pro Gln Phe Arg Glu Leu
66   210                              215                              220
67   Gly Tyr Val Leu Lys Asp Lys Tyr Asn Glu Ala Val His Val Glu Pro
68   225                              230                              235                              240
69   Val Arg Ala Ser Arg Asn Lys Arg Pro Thr Phe Leu Lys Ile Lys Lys
70                               245                               250                               255
71   Pro Leu Ser Tyr Arg Lys Pro Met Asp Thr Asp Leu Val Tyr Ile Glu
72                               260                               265                               270
73   Lys Ser Pro Asn Tyr Cys Glu Glu Asp Pro Val Thr Gly Ser Val Gly
74   275                              280                              285
75   Thr Gln Gly Arg Ala Cys Asn Lys Thr Ala Pro Gln Ala Ser Gly Cys
76   290                              295                              300
77   Asp Leu Met Cys Cys Gly Arg Gly Tyr Asn Thr His Gln Tyr Ala Arg
78   305                              310                              315                              320
79   Val Trp Gln Cys Asn Cys Lys Phe His Trp Cys Cys Tyr Val Lys Cys
80                               325                               330                               335
81   Asn Thr Cys Ser Glu Arg Thr Glu Met Tyr Thr Cys Lys
82                               340                               345

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VERIFICATION SUMMARY
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Line ? Error/Warning

Original Text
